RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/651,668A
Source: IFW16
Date Processed by STIC: 07/10/2006

ENTERED

IFW16

RAW SEQUENCE LISTING DATE: 07/10/2006
PATENT APPLICATION: **US/10/651,668A** TIME: 12:22:45

Input Set : **F:\ISPA-5001-C1 (Substitute Seq List).ST25.txt**
Output Set: **N:\CRF4\07102006\J651668A.raw**

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 3 <110> APPLICANT: Takeda San Diego, Inc.
 5 <120> TITLE OF INVENTION: CRYSTALLIZATION OF ISPA
 7 <130> FILE REFERENCE: SYR-IspA-5001-C1
 9 <140> CURRENT APPLICATION NUMBER: 10/651,668A
10 <141> CURRENT FILING DATE: 2003-08-28
12 <160> NUMBER OF SEQ ID NOS: 2
14 <170> SOFTWARE: PatentIn version 3.3
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 314
18 <212> TYPE: PRT
19 <213> ORGANISM: Artificial
21 <220> FEATURE:
22 <223> OTHER INFORMATION: Amino acid sequence for full-length E. coli IspA with an
23       N-terminal His-tag
26 <220> FEATURE:
27 <221> NAME/KEY: MISC_FEATURE
28 <222> LOCATION: (1)..(15)
29 <223> OTHER INFORMATION: N-terminal His-tag
31 <220> FEATURE:
32 <221> NAME/KEY: MISC_FEATURE
33 <222> LOCATION: (16)..(314)
34 <223> OTHER INFORMATION: Full-length E. coli IspA
36 <400> SEQUENCE: 1
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39 1               5                  10                  15
42 Asp Phe Pro Gln Gln Leu Glu Ala Cys Val Lys Gln Ala Asn Gln Ala
43             20                  25                  30
46 Leu Ser Arg Phe Ile Ala Pro Leu Pro Phe Gln Asn Thr Pro Val Val
47         35                  40                  45
50 Glu Thr Met Gln Tyr Gly Ala Leu Leu Gly Gly Lys Arg Leu Arg Pro
51     50                  55                  60
54 Phe Leu Val Tyr Ala Thr Gly His Met Phe Gly Val Ser Thr Asn Thr
55 65                  70                  75                  80
58 Leu Asp Ala Pro Ala Ala Ala Val Glu Cys Ile His Ala Tyr Ser Leu
59                 85                  90                  95
62 Ile His Asp Asp Leu Pro Ala Met Asp Asp Asp Asp Leu Arg Arg Gly
63             100                 105                 110
66 Leu Pro Thr Cys His Val Lys Phe Gly Glu Ala Asn Ala Ile Leu Ala
67         115                 120                 125
70 Gly Asp Ala Leu Gln Thr Leu Ala Phe Ser Ile Leu Ser Asp Ala Asp
71     130                 135                 140
74 Met Pro Glu Val Ser Asp Arg Asp Arg Ile Ser Met Ile Ser Glu Leu
75 145                 150                 155                 160
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78 Ala Ser Ala Ser Gly Ile Ala Gly Met Cys Gly Gly Gln Ala Leu Asp
79                 165                 170                 175
82 Leu Asp Ala Glu Gly Lys His Val Pro Leu Asp Ala Leu Glu Arg Ile
83             180                 185                 190
86 His Arg His Lys Thr Gly Ala Leu Ile Arg Ala Ala Val Arg Leu Gly
87         195                 200                 205
90 Ala Leu Ser Ala Gly Asp Lys Gly Arg Arg Ala Leu Pro Val Leu Asp
91     210                 215                 220
94 Lys Tyr Ala Glu Ser Ile Gly Leu Ala Phe Gln Val Gln Asp Asp Ile
95 225                 230                 235                 240
98 Leu Asp Val Val Gly Asp Thr Ala Thr Leu Gly Lys Arg Gln Gly Ala
99                 245                 250                 255
102 Asp Gln Gln Leu Gly Lys Ser Thr Tyr Pro Ala Leu Leu Gly Leu Glu
103             260                 265                 270
106 Gln Ala Arg Lys Lys Ala Arg Asp Leu Ile Asp Asp Ala Arg Gln Ser
107         275                 280                 285
110 Leu Lys Gln Leu Ala Glu Gln Ser Leu Asp Thr Ser Ala Leu Glu Ala
111     290                 295                 300
114 Leu Ala Asp Tyr Ile Ile Gln Arg Asn Lys
115 305                 310
118 <210> SEQ ID NO: 2
119 <211> LENGTH: 945
120 <212> TYPE: DNA
121 <213> ORGANISM: Artificial
123 <220> FEATURE:
124 <223> OTHER INFORMATION: cDNA sequence encoding IspA with an N-terminal His-tag
127 <220> FEATURE:
128 <221> NAME/KEY: misc_feature
129 <222> LOCATION: (1)..(45)
130 <223> OTHER INFORMATION: Sequence encoding N-terminal His-tag
132 <220> FEATURE:
133 <221> NAME/KEY: misc_feature
134 <222> LOCATION: (46)..(945)
135 <223> OTHER INFORMATION: Sequence encoding full-length E. coli IspA
137 <400> SEQUENCE: 2
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140 caactcgaag cctgcgttaa gcaggccaac caggcgctga gccgttttat cgccccactg    120
142 ccctttcaga acactcccgt ggtcgaaacc atgcagtatg gcgcattatt aggtggtaag    180
144 cgcctgcgac ctttcctggt ttatgccacc ggtcatatgt tcggcgttag cacaaacacg    240
146 ctggacgcac ccgctgccgc cgttgagtgt atccacgctt actcattaat tcatgatgat    300
148 ttaccggcaa tggatgatga cgatctgcgt cgcggtttgc caacctgcca tgtgaagttt    360
150 ggcgaagcaa acgcgattct cgctggcgac gctttacaaa cgctggcgtt ctcgatttta    420
152 agcgatgccg atatgccgga agtgtcggac cgcgacagaa tttcgatgat ttctgaactg    480
154 gcgagcgcca gtggtattgc cggaatgtgc ggtggtcagg cattagattt agacgcggaa    540
156 ggcaaacacg tacctctgga cgcgcttgag cgtattcatc gtcataaaac cggcgcattg    600
158 attcgcgccg ccgttcgcct tggtgcatta agcgccggag ataaaggacg tcgtgctctg    660
160 ccggtactcg acaagtatgc agagagcatc ggccttgcct tccaggttca ggatgacatc    720
162 ctggatgtgg tgggagatac tgcaacgttg ggaaaacgcc agggtgccga ccagcaactt    780
164 ggtaaaagta cctaccctgc acttctgggt cttgagcaag cccggaagaa agcccgggat    840
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166 ctgatcgacg atgcccgtca gtcgctgaaa caactggctg aacagtcact cgatacctcg 900
168 gcactggaag cgctagcgga ctacatcatc cagcgtaata aataa 945

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/10/2006
PATENT APPLICATION: **US/10/651,668A** TIME: 12:22:46

Input Set : **F:\ISPA-5001-C1 (Substitute Seq List).ST25.txt**
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Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2